

PCT10

DATE: 03/11/2002 RAW SEQUENCE LISTING PATENT APPLICATION: US/10/069,385 TIME: 10:39:26 Input Set : A:\X-13268.App.txt mai Output Set: N:\CRF3\03112002\J069385.raw Does North State 4 <110> APPLICANT: Witcher, Derrick C'orrer's de dans 5 Tian, Yu 6 Atkinson, Paul 8 <120> TITLE OF INVENTION: FLINT Analog Compounds and Formulations Thereof 10 <130> FILE REFERENCE: X-13268 CHS 12 <140> CURRENT AFFILIATE: 2002-02-19 13 <141> CURRENT FILING DATE: 2002-02-19 > 12 <140> CURRENT APPLICATION NUMBER: US/10/069,385 15 <160> NUMBER OF SEQ ID NOS: 3

17 <170> SOFTWARE: PatentIn Ver. 2.0

## ERRORED SEQUENCES

 $77 \ll 210$ > SEQ ID NO: 2 78 - 211> LENGTH: 300 79 <212> TYPE: PRT 80 <213> ORGANISM: Homo sapiens E - -> 82 < 400 > SEQUENCE: (3) 283 Met Arg Ala Leu Ĝlu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val Leu 84 1 5 10 15 86 Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly Val Ala Glu 87 20 25 30 89 Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val 90 35 45 40 92 Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg 93 50 55 60 95 Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln 96 65 70 75 80 98 Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly 94 90 85 95 101 Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His Asn Arg Ala 102 100 105 110 104 Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu 105 120 115 125 107 His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro 135 108130 140 110 Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala 111 145 150 155 160113 Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala 114 165 170 175 116 Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser His Asp Thr Leu 117 185 180 190 119 Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala

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122 Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile 125 Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu 126 225 128 Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys 1.29 131 Lou Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu 134 Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu 137 Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 141 <210> SEQ ID NO: 3 142 <211> LENGTH: 936 143 <212> TYPE: DNA 144 - 213> ORGANISM: Homo sapiens 146 - 220> FEATURE:  $147 \rightarrow 221$  NAME/KEY: CDS  $148 \ll 222 > LOCATION: (25)..(924)$ 150 <400> SEQUENCE: 3 151 geteteeetg eteeageaag gaee atg agg geg etg gag ggg eea gge etg Met Arg Ala Leu Glu Gly Pro Gly Leu 156 Ser Leu Leu Cys Leu Val Leu Ala Leu Pro Ala Leu Leu Pro Val Pro 159 get gta ege gga gtg gea gaa aca eee ace tae eee tgg egg gae gea 160 Ala Val Arg Gly Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala 163 gag aca ggg gag egg etg gtg tge gee eag tge eee eea gge ace tit 164 Glu Thr Gly Glu Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe 167 gtg cag cgg ccg tgc cgc cga gac age ccc acg acg tgt ggc ccg tgt 168 Val Gln Arg Pro Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys 171 dea deg ege eac tac acg eag the tgg aac tac etg gag ege tge ege 172 Pro Pro Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg 175 tae tge aae gte ete tge ggg gag egt gag gag gag gea egg get tge 176 Tyr Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys 179 cae gee ace cae aac egt gee tge ege tge ege ace gge tte tte geg 180 His Ala Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala 183 cac get ggt the tige fig gag cae gea teg tigt eea eet ggt gee gge 184 His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly 187 gtg att gee eeg gge ace eee age eag aae aeg eag tge eag eeg tge 

RAW SEQUENCE LISTINGDAPATENT APPLICATION:US/10/069,385T1

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188 Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys 191 dee dea gge ace the tea gee age age tee age tea gag eag tge eag 142 Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser Glu Gln Cys Gln 195 dee dae dige aad tige acg gee dig gige dig gee die att gig dea gige 196 Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala Leu Ile Val Pro Gly 197 170 199 tot too too cat gad add ofg fige acc age fige act gge the eee etc. 200 Ser Ser Ser His Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu age ace agg gta eea gga get gag gag tgt gag egt gee gte ate gae 204 Ser Thr Arg Val Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp tit gig get the cag gae ale tee ale aag agg eig eag egg eig eig 208 Phe Val Ala Phe Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu 211 cag gee ete gag gee eeg gag gge tgg get eeg aca eea agg geg gge 212 Gln Ala Leu Glu Ala Pro Glu Gly Trp Ala Pro Thr Pro Arg Ala Gly ege geg gee tig eag eig aag eig egi egg eig ace aeg gag eie eig 216 Arg Ala Ala Leu Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu 217 250 ggg geg cag gac ggg geg etg etg gtg egg etg etg eag geg etg ege 220 Gly Ala Gln Asp Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg gtg gee agg atg eee ggg etg gag egg age gte egt gag ege tte etc 224 Val Ala Arg Met Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu 227 cet gtg cae tgateetgge ee 228 Pro Val His E--> 237/ X-13268 E--> 245(1

VERIFICATION SUMMARY PATENT APPLICATION: US/10/069,385 DATE: 03/11/2002 TIME: 10:39:27

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L:12 M:270 C: Current Application Number differs, Replaced Application Number L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:82 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:3 L:237 M:254 E: No. of Bases conflict, LENGTH:Input:-13268 Counted:938 SEQ:3 L:237 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1 M:254 Repeated in SeqNo=3 L:245 M:252 E: No. of Seq. differs, <211>LENGTH:Input:936 Found:938 SEQ:3